## What is claimed is:

- 1. An isolated polypeptide comprising an amino acid sequence selected from the group consisting of:
  - a) a mature form of the amino acid sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14;
  - b) a variant of a mature form of the amino acid sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14, wherein any amino acid in the mature form is changed to a different amino acid, provided that no more than 15% of the amino acid residues in the sequence of the mature form are so changed;
  - the amino acid sequence selected from the group consisting of SEQ ID NO:
    8, 10, 12 and 14;
  - d) a variant of the amino acid sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14, wherein any amino acid specified in the chosen sequence is changed to a different amino acid, provided that no more than 15% of the amino acid residues in the sequence are so changed; and
  - e) a fragment of any of a) through d).
- 2. The polypeptide of claim 1 that is a naturally occurring allelic variant of the sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14.
- 3. The polypeptide of claim 2, wherein the variant is the translation of a single nucleotide polymorphism.
- 4. The polypeptide of claim 1 that is a variant polypeptide described therein, wherein any amino acid specified in the chosen sequence is changed to provide a conservative substitution.
- 5. An isolated nucleic acid molecule comprising a nucleic acid sequence encoding a polypeptide comprising an amino acid sequence selected from the group consisting of:
  - a) a mature form of the amino acid sequence given SEQ ID NO: 8, 10, 12 and 14:
  - b) a variant of a mature form of the amino acid sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14, wherein any amino acid

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- in the mature form of the chosen sequence is changed to a different amino acid, provided that no more than 15% of the amino acid residues in the sequence of the mature form are so changed;
- the amino acid sequence selected from the group consisting of SEQ ID NO:
   8, 10, 12 and 14;
- d) a variant of the amino acid sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14, in which any amino acid specified in the chosen sequence is changed to a different amino acid, provided that no more than 15% of the amino acid residues in the sequence are so changed;
- e) a nucleic acid fragment encoding at least a portion of a polypeptide comprising the amino acid sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14, or any variant of said polypeptide wherein any amino acid of the chosen sequence is changed to a different amino acid, provided that no more than 10% of the amino acid residues in the sequence are so changed; and
- f) the complement of any of said nucleic acid molecules.
- 6. The nucleic acid molecule of claim 5, wherein the nucleic acid molecule comprises the nucleotide sequence of a naturally occurring allelic nucleic acid variant.
- 7. The nucleic acid molecule of claim 5 that encodes a variant polypeptide, wherein the variant polypeptide has the polypeptide sequence of a naturally occurring polypeptide variant.
- 8. The nucleic acid molecule of claim 5, wherein the nucleic acid molecule comprises a single nucleotide polymorphism encoding said variant polypeptide.
- 9. The nucleic acid molecule of claim 5, wherein said nucleic acid molecule comprises a nucleotide sequence selected from the group consisting of
  - a) the nucleotide sequence selected from the group consisting of SEQ ID NO: 7,9, 11 and 13;
  - b) a nucleotide sequence wherein one or more nucleotides in the nucleotide sequence selected from the group consisting of SEQ ID NO: 7, 9, 11 and 13 is changed from that selected from the group consisting of the chosen sequence to

- a different nucleotide provided that no more than 15% of the nucleotides are so changed;
- c) a nucleic acid fragment of the sequence selected from the group consisting of SEQ ID NO: 7, 9, 11 and 13; and
- d) a nucleic acid fragment wherein one or more nucleotides in the nucleotide sequence selected from the group consisting of SEQ ID NO: 7, 9, 11 and 13 is changed from that selected from the group consisting of the chosen sequence to a different nucleotide provided that no more than 15% of the nucleotides are so changed.
- The nucleic acid molecule of claim 5, wherein said nucleic acid molecule hybridizes under stringent conditions to the nucleotide sequence selected from the group consisting of SEQ ID NO: 7, 9, 11 and 13, or a complement of said nucleotide sequence.
- 11. The nucleic acid molecule of claim 5, wherein the nucleic acid molecule comprises a nucleotide sequence in which any nucleotide specified in the coding sequence of the chosen nucleotide sequence is changed from that selected from the group consisting of the chosen sequence to a different nucleotide provided that no more than 15% of the nucleotides in the chosen coding sequence are so changed, an isolated second polynucleotide that is a complement of the first polynucleotide, or a fragment of any of them.
- 12. A vector comprising the nucleic acid molecule of claim 11.
- 13. The vector of claim 12, further comprising a promoter operably linked to said nucleic acid molecule.
- 14. A cell comprising the vector of claim 12.
- 15. An antibody that binds immunospecifically to the polypeptide of claim 1.
- 16. The antibody of claim 15, wherein said antibody is a monoclonal antibody.
- 17. The antibody of claim 15, wherein the antibody is a humanized antibody.

- 18. A method for determining the presence or amount of the polypeptide of claim 1 in a sample, the method comprising:
  - (a) providing said sample;
  - (b) introducing said sample to an antibody that binds immunospecifically to the polypeptide; and
  - (c) determining the presence or amount of antibody bound to said polypeptide, thereby determining the presence or amount of polypeptide in said sample.
- 19. A method for modulating the activity of the polypeptide of claim 1, the method comprising introducing a cell sample expressing the polypeptide of said claim with a compound that binds to said polypeptide in an amount sufficient to modulate the activity of the polypeptide.
- 20. A method of treating or preventing a pathology associated with the polypeptide of claim 1, said method comprising administering the polypeptide of claim 1 to a subject in which such treatment or prevention is desired in an amount sufficient to treat or prevent said pathology in said subject.
- 21. The method of claim 20, wherein said subject is a human.
- 22. A method of treating or preventing a pathology associated with the polypeptide of claim 1, said method comprising administering to a subject in which such treatment or prevention is desired a nucleic acid selected from the group consisting of SEQ ID NO: 7, 9, 11 and 13, in an amount sufficient to treat or prevent said pathology in said subject.
- 23. The method of claim 22, wherein said subject is a human.
- A method of treating or preventing a pathology associated with the polypeptide of claim 1, said method comprising administering to a subject in which such treatment or prevention is desired an antibody selected from the group consisting of an antibody to SEQ ID NO:
  8, 10, 12 and 14, in an amount sufficient to treat or prevent said pathology in said subject.
- 25. The method of claim 24, wherein the subject is a human.

- 26. A pharmaceutical composition comprising the polypeptide of claim 1 and a pharmaceutically acceptable carrier.
- 27. A pharmaceutical composition comprising the nucleic acid molecule of claim 5 and a pharmaceutically acceptable carrier.
- 28. A pharmaceutical composition comprising the antibody of claim 15 and a pharmaceutically acceptable carrier.
- 29. A kit comprising in one or more containers, the pharmaceutical composition of claim 26.
- 30. A kit comprising in one or more containers, the pharmaceutical composition of claim 27.
- 31. A kit comprising in one or more containers, the pharmaceutical composition of claim 28.
- 32. A method for screening for a modulator of activity or of latency or predisposition to a pathology associated with the polypeptide of claim 1, said method comprising:
  - a. administering a test compound to a test animal at increased risk for a pathology associated with the polypeptide of claim 1, wherein said test animal recombinantly expresses the polypeptide of claim 1;
  - b. measuring the activity of said polypeptide in said test animal after administering the compound of step (a); and
  - c. comparing the activity of said protein in said test animal with the activity of said polypeptide in a control animal not administered said polypeptide, wherein a change in the activity of said polypeptide in said test animal relative to said control animal indicates the test compound is a modulator of latency of, or predisposition to, a pathology associated with the polypeptide of claim 1.

d.

- 33. The method of claim 32, wherein said test animal is a recombinant test animal that expresses a test protein transgene or expresses said transgene under the control of a promoter at an increased level relative to a wild-type test animal, and wherein said promoter is not the native gene promoter of said transgene.
- 34. A method of treating a pathological state in a mammal, the method comprising administering to the mammal a polypeptide in an amount that is sufficient to alleviate the

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pathological state, wherein the polypeptide is a polypeptide having an amino acid sequence at least 95% identical to a polypeptide comprising the amino acid sequence selected from the group consisting of SEQ ID NO: 8, 10, 12 and 14, or a biologically active fragment thereof.

35. A method of treating a pathological state in a mammal, the method comprising administering to the mammal the antibody of claim 15 in an amount sufficient to alleviate the pathological state.